STIC Biotechnology Systems Branch

RAW SEQUENCE LISTING ERROR REPORT

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following computer readable form:

Application Serial Number: 10/575,261
Source: IFW P
Date Processed by STIC: 04/24/2006

THE ATTACHED PRINTOUT EXPLAINS DETECTED ERRORS.
PLEASE FORWARD THIS INFORMATION TO THE APPLICANT BY EITHER:
1) **INCLUDING A COPY OF THIS PRINTOUT IN YOUR NEXT COMMUNICATION TO THE APPLICANT, WITH A NOTICE TO COMPLY or,**
2) **TELEPHONING APPLICANT AND FAXING A COPY OF THIS PRINTOUT, WITH A NOTICE TO COMPLY**

FOR CRF SUBMISSION AND PATENTIN SOFTWARE QUESTIONS, PLEASE CONTACT MARK SPENCER, TELEPHONE: 571-272-2510; FAX: 571-273-0221

TO **REDUCE** ERRORED SEQUENCE LISTINGS, **PLEASE** USE THE **CHECKER VERSION 4.4.0 PROGRAM**, ACCESSIBLE THROUGH THE U.S. PATENT AND TRADEMARK OFFICE WEBSITE. SEE BELOW FOR ADDRESS:
http://www.uspto.gov/web/offices/pac/checker/chkrnote.htm

Applicants submitting genetic sequence information electronically on diskette or CD-Rom should be aware that there is a possibility that the disk/CD-Rom may have been affected by treatment given to all incoming mail.
Please consider using alternate methods of submission for the disk/CD-Rom or replacement disk/CD-Rom.
Any reply including a sequence listing in electronic form should NOT be sent to the 20231 zip code address for the United States Patent and Trademark Office, and instead should be sent via the following to the indicated addresses:

1. EFS-Bio (<**http://www.uspto.gov/ebc/efs/downloads/documents.htm**> , **EFS Submission User Manual** - ePAVE)
2. **U.S. Postal Service:** Commissioner for Patents, P.O. Box 1450, Alexandria, VA 22313-1450
3. **Hand Carry, Federal Express, United Parcel Service, or other delivery service (EFFECTIVE 01/14/05):** U.S. Patent and Trademark Office, Mail Stop Sequence, Customer Window, Randolph Building, 401 Dulany Street, Alexandria, VA 22314

Revised 01/10/06

IFWP

RAW SEQUENCE LISTING DATE: 04/24/2006
PATENT APPLICATION: **US/10/575,261** TIME: 16:15:41

Input Set : **A:\seq list.txt**
Output Set: **N:\CRF4\04242006\J575261.raw**

```
     3 <110> APPLICANT: KYOWA HAKKO KOGYO CO., LTD.
     5 <120> TITLE OF INVENTION: Fusion protein composition
     7 <130> FILE REFERENCE: 11613WO1
C--> 9 <140> CURRENT APPLICATION NUMBER: US/10/575,261
C--> 9 <141> CURRENT FILING DATE: 2006-04-10
     9 <150> PRIOR APPLICATION NUMBER: P2003-350158
    10 <151> PRIOR FILING DATE: 2003-10-08
    12 <160> NUMBER OF SEQ ID NOS: 113
    14 <170> SOFTWARE: PatentIn Ver. 2.1
```

Does Not Comply
Corrected Diskette Needed
(CRF-1, 2, 4, 5, 6)

ERRORED SEQUENCES

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   131 <210> SEQ ID NO: 2
   132 <211> LENGTH: 376 -> found 372
   133 <212> TYPE: PRT
   134 <213> ORGANISM: Cricetulus griseus
   136 <400> SEQUENCE: 2
   137 Met Ala His Ala Pro Ala Ser Cys Pro Ser Ser Arg Asn Ser Gly Asp
   138   1               5                  10                  15
   140 Gly Asp Lys Gly Lys Pro Arg Lys Val Ala Leu Ile Thr Gly Ile Thr
   141              20                  25                  30
   143 Gly Gln Asp Gly Ser Tyr Leu Ala Glu Phe Leu Leu Glu Lys Gly Tyr
   144          35                  40                  45
   146 Glu Val His Gly Ile Val Arg Arg Ser Ser Ser Phe Asn Thr Gly Arg
   147      50                  55                  60
   149 Ile Glu His Leu Tyr Lys Asn Pro Gln Ala His Ile Glu Gly Asn Met
   150  65                  70                  75                  80
   152 Lys Leu His Tyr Gly Asp Leu Thr Asp Ser Thr Cys Leu Val Lys Ile
E--> 153 85              85  90              90  95              95 100
   155 Ile Asn Glu Val Lys Pro Thr Glu Ile Tyr Asn Leu Gly Ala Gln Ser
E--> 156             100 105             105 110             110 115
   158 His Val Lys Ile Ser Phe Asp Leu Ala Glu Tyr Thr Ala Asp Val Asp
E--> 159         115 120             120 125             125 130
   161 Gly Val Gly Thr Leu Arg Leu Leu Asp Ala Ile Lys Thr Cys Gly Leu
E--> 162     130 135             135 140             140 145
   164 Ile Asn Ser Val Lys Phe Tyr Gln Ala Ser Thr Ser Glu Leu Tyr Gly
E--> 165 145 150             150 155             155 160             160
   167 Lys Val Gln Glu Ile Pro Gln Lys Glu Thr Thr Pro Phe Tyr Pro Arg
E--> 168 165             165 170             170 175             175 180
   170 Ser Pro Tyr Gly Ala Ala Lys Leu Tyr Ala Tyr Trp Ile Val Val Asn
E--> 171             180 185             185 190             190 195
   173 Phe Arg Glu Ala Tyr Asn Leu Phe Ala Val Asn Gly Ile Leu Phe Asn
                195                 200                 205
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Invalid Amino Acid numbering

RAW SEQUENCE LISTING DATE: 04/24/2006
PATENT APPLICATION: US/10/575,261 TIME: 16:15:41

Input Set : A:\seq list.txt
Output Set: N:\CRF4\04242006\J575261.raw

```
E--> 174                200                 205                 210
     176 His Glu Ser Pro Arg Arg Gly Ala Asn Phe Val Thr Arg Lys Ile Ser
E--> 177      210 215              215 220              220 225
     179 Arg Ser Val Ala Lys Ile Tyr Leu Gly Gln Leu Glu Cys Phe Ser Leu
E--> 180 225 230              230 235              235 240              240
     182 Gly Asn Leu Asp Ala Lys Arg Asp Trp Gly His Ala Lys Asp Tyr Val
E--> 183 245              245 250              250 255              255 260
     185 Glu Ala Met Trp Leu Met Leu Gln Asn Asp Glu Pro Glu Asp Phe Val
E--> 186              260 265              265 270              270 275
     188 Ile Ala Thr Gly Glu Val His Ser Val Arg Glu Phe Val Glu Lys Ser
E--> 189         245 280              280 285              285 290
     191 Phe Met His Ile Gly Lys Thr Ile Val Trp Glu Gly Lys Asn Glu Asn
E--> 192      290 295              295 300              300 305
     194 Glu Val Gly Arg Cys Lys Glu Thr Gly Lys Ile His Val Thr Val Asp
E--> 195 305 310              310 315              315 320              320
     197 Leu Lys Tyr Tyr Arg Pro Thr Glu Val Asp Phe Leu Gln Gly Asp Cys
E--> 198 325              325 330              330 335              335 340
     200 Ser Lys Ala Gln Gln Lys Leu Asn Trp Lys Pro Arg Val Ala Phe Asp
E--> 201              240 345              245 350              250 355
     204 Glu Leu Val Arg Glu Met Val Gln Ala Asp Val Glu Leu Met Arg Thr
E--> 205         355 360              260 365              265 370
     207 Asn Pro Asn Ala
E--> 208      370 375
     471 <210> SEQ ID NO: 7
     472 <211> LENGTH: 575
     473 <212> TYPE: PRT
     474 <213> ORGANISM: Cricetulus griseus
     476 <400> SEQUENCE: 7
     477 Met Arg Ala Trp Thr Gly Ser Trp Arg Trp Ile Met Leu Ile Leu Phe
     478   1               5                  10                  15
     480 Ala Trp Gly Thr Leu Leu Phe Tyr Ile Gly Gly His Leu Val Arg Asp
     481              20                  25                  30
     483 Asn Asp His Pro Asp His Ser Ser Arg Glu Leu Ser Lys Ile Leu Ala
     484          35                  40                  45
     486 Lys Leu Glu Arg Leu Lys Gln Gln Asn Glu Asp Leu Arg Arg Met Ala
     487      50                  55                  60
     489 Glu Ser Leu Arg Ile Pro Glu Gly Pro Ile Asp Gln Gly Thr Ala Thr
     490  65                  70                  75                  80
     492 Gly Arg Val Arg Val Leu Glu Glu Gln Leu Val Lys Ala Lys Glu Gln
     493                  85                  90                  95
     495 Ile Glu Asn Tyr Lys Lys Gln Ala Arg Asn Asp Leu Gly Lys Asp His
     496              100                 105                 110
     498 Glu Ile Leu Arg Arg Arg Ile Glu Asn Gly Ala Lys Glu Leu Trp Phe
     499          115                 120                 125
     501 Phe Leu Gln Ser Glu Leu Lys Lys Leu Lys Lys Leu Glu Gly Asn Glu
     502      130                 135                 140
     505 Leu Gln Arg His Ala Asp Glu Ile Leu Leu Asp Leu Gly His His Glu
     506 145                 150                 155                 160
     508 Arg Ser Ile Met Thr Asp Leu Tyr Tyr Leu Ser Gln Thr Asp Gly Ala
```

Same Error

RAW SEQUENCE LISTING DATE: 04/24/2006
PATENT APPLICATION: US/10/575,261 TIME: 16:15:41

Input Set : A:\seq list.txt
Output Set: N:\CRF4\04242006\J575261.raw

```
509                 165                 170                 175
511 Gly Glu Trp Arg Glu Lys Glu Ala Lys Asp Leu Thr Glu Leu Val Gln
512             180                 185                 190
514 Arg Arg Ile Thr Tyr Leu Gln Asn Pro Lys Asp Cys Ser Lys Ala Arg
515         195                 200                 205
517 Lys Leu Val Cys Asn Ile Asn Lys Gly Cys Gly Tyr Gly Cys Gln Leu
518     210                 215                 220
520 His His Val Val Tyr Cys Phe Met Ile Ala Tyr Gly Thr Gln Arg Thr
521 225                 230                 235                 240
523 Leu Ile Leu Glu Ser Gln Asn Trp Arg Tyr Ala Thr Gly Gly Trp Glu
524                 245                 250                 255
526 Thr Val Phe Arg Pro Val Ser Glu Thr Cys Thr Asp Arg Ser Gly Leu
527             260                 265                 270
529 Ser Thr Gly His Trp Ser Gly Glu Val Lys Asp Lys Asn Val Gln Val
530         275                 280                 285
532 Val Glu Leu Pro Ile Val Asp Ser Leu His Pro Arg Pro Pro Tyr Leu
533     290                 295                 300
535 Pro Leu Ala Val Pro Glu Asp Leu Ala Asp Arg Leu Leu Arg Val His
536 305                 310                 315                 320
538 Gly Asp Pro Ala Val Trp Trp Val Ser Gln Phe Val Lys Tyr Leu Ile
539                 325                 330                 335
541 Arg Pro Gln Pro Trp Leu Glu Arg Glu Ile Glu Glu Thr Thr Lys Lys
542             340                 345                 350
544 Leu Gly Phe Lys His Pro Val Ile Gly Val His Val Arg Arg Thr Asp
545         355                 360                 365
547 Lys Val Gly Thr Glu Ala Ala Phe His Pro Ile Glu Glu Tyr Met Val
548     370                 375                 380
550 His Val Glu Glu His Phe Gln Leu Leu Glu Arg Arg Met Lys Val Asp
551 385                 390                 395                 400
553 Lys Lys Arg Val Tyr Leu Ala Thr Asp Asp Pro Ser Leu Leu Lys Glu
554                 405                 410                 415
556 Ala Lys Thr Lys Tyr Ser Asn Tyr Glu Phe Ile Ser Asp Asn Ser Ile
557             420                 425                 430
559 Ser Trp Ser Ala Gly Leu His Asn Arg Tyr Thr Glu Asn Ser Leu Arg
560         435                 440                 445
562 Gly Val Ile Leu Asp Ile His Phe Leu Ser Gln Ala Asp Phe Leu Val
563     450                 455                 460
565 Cys Thr Phe Ser Ser Gln Val Cys Arg Val Ala Tyr Glu Ile Met Gln
566 465                 470                 475                 480
568 Thr Leu His Pro Asp Ala Ser Ala Asn Phe His Ser Leu Asp Asp Ile
569                 485                 490                 495
571 Tyr Tyr Phe Gly Gly Gln Asn Ala His Asn Gln Ile Ala Val Tyr Pro
572             500                 505                 510
574 His Gln Pro Arg Thr Lys Glu Glu Ile Pro Met Glu Pro Gly Asp Ile
575         515                 520                 525
577 Ile Gly Val Ala Gly Asn His Trp Asn Gly Tyr Ser Lys Gly Val Asn
578     530                 535                 540
580 Arg Lys Leu Gly Lys Thr Gly Leu Tyr Pro Ser Tyr Lys Val Arg Glu
581 545                 550                 555                 560
```

RAW SEQUENCE LISTING DATE: 04/24/2006
PATENT APPLICATION: **US/10/575,261** TIME: 16:15:41

Input Set : **A:\seq list.txt**
Output Set: **N:\CRF4\04242006\J575261.raw**

```
     583 Lys Ile Glu Thr Val Lys Tyr Pro Thr Tyr Pro Glu Ala Glu Lys
E--> 584                 565                 570                 575
     587 <210> SEQ ID NO: 8
     588 <211> LENGTH: 575
     589 <212> TYPE: PRT
     590 <213> ORGANISM: Mus musculus
     592 <400> SEQUENCE: 8
     593 Met Arg Ala Trp Thr Gly Ser Trp Arg Trp Ile Met Leu Ile Leu Phe
     594   1               5                  10                  15
     596 Ala Trp Gly Thr Leu Leu Phe Tyr Ile Gly Gly His Leu Val Arg Asp
     597              20                  25                  30
     599 Asn Asp His Pro Asp His Ser Ser Arg Glu Leu Ser Lys Ile Leu Ala
     600          35                  40                  45
     602 Lys Leu Glu Arg Leu Lys Gln Gln Asn Glu Asp Leu Arg Arg Met Ala
     604      50                  55                  60
     606 Glu Ser Leu Arg Ile Pro Glu Gly Pro Ile Asp Gln Gly Thr Ala Thr
     607  65                  70                  75                  80
     609 Gly Arg Val Arg Val Leu Glu Glu Gln Leu Val Lys Ala Lys Glu Gln
     610                  85                  90                  95
     612 Ile Glu Asn Tyr Lys Lys Gln Ala Arg Asn Gly Leu Gly Lys Asp His
     613             100                 105                 110
     615 Glu Ile Leu Arg Arg Arg Ile Glu Asn Gly Ala Lys Glu Leu Trp Phe
     616         115                 120                 125
     618 Phe Leu Gln Ser Glu Leu Lys Lys Leu Lys His Leu Glu Gly Asn Glu
     619     130                 135                 140
     621 Leu Gln Arg His Ala Asp Glu Ile Leu Leu Asp Leu Gly His His Glu
     622 145                 150                 155                 160
     624 Arg Ser Ile Met Thr Asp Leu Tyr Tyr Leu Ser Gln Thr Asp Gly Ala
     625                 165                 170                 175
     627 Gly Asp Trp Arg Glu Lys Glu Ala Lys Asp Leu Thr Glu Leu Val Gln
     628             180                 185                 190
     630 Arg Arg Ile Thr Tyr Leu Gln Asn Pro Lys Asp Cys Ser Lys Ala Arg
     631         195                 200                 205
     633 Lys Leu Val Cys Asn Ile Asn Lys Gly Cys Gly Tyr Gly Cys Gln Leu
     634     210                 215                 220
     636 His His Val Val Tyr Cys Phe Met Ile Ala Tyr Gly Thr Gln Arg Thr
     637 225                 230                 235                 240
     639 Leu Ile Leu Glu Ser Gln Asn Trp Arg Tyr Ala Thr Gly Gly Trp Glu
     640                 245                 250                 255
     642 Thr Val Phe Arg Pro Val Ser Glu Thr Cys Thr Asp Arg Ser Gly Leu
     643             260                 265                 270
     645 Ser Thr Gly His Trp Ser Gly Glu Val Asn Asp Lys Asn Ile Gln Val
     646         275                 280                 285
     648 Val Glu Leu Pro Ile Val Asp Ser Leu His Pro Arg Pro Pro Tyr Leu
     649     290                 295                 300
     651 Pro Leu Ala Val Pro Glu Asp Leu Ala Asp Arg Leu Leu Arg Val His
     652 305                 310                 315                 320
     654 Gly Asp Pro Ala Val Trp Trp Val Ser Gln Phe Val Lys Tyr Leu Ile
     655                 325                 330                 335
```

RAW SEQUENCE LISTING DATE: 04/24/2006
PATENT APPLICATION: US/10/575,261 TIME: 16:15:41

Input Set : A:\seq list.txt
Output Set: N:\CRF4\04242006\J575261.raw

```
657 Arg Pro Gln Pro Trp Leu Glu Lys Glu Ile Glu Glu Ala Thr Lys Lys
658             340                 345                 350
660 Leu Gly Phe Lys His Pro Val Ile Gly Val His Val Arg Arg Thr Asp
661         355                 360                 365
663 Lys Val Gly Thr Glu Ala Ala Phe His Pro Ile Glu Glu Tyr Met Val
664     370                 375                 380
666 His Val Glu Glu His Phe Gln Leu Leu Ala Arg Arg Met Gln Val Asp
667 385                 390                 395                 400
669 Lys Lys Arg Val Tyr Leu Ala Thr Asp Asp Pro Thr Leu Leu Lys Glu
670                 405                 410                 415
672 Ala Lys Thr Lys Tyr Ser Asn Tyr Glu Phe Ile Ser Asp Asn Ser Ile
673             420                 425                 430
675 Ser Trp Ser Ala Gly Leu His Asn Arg Tyr Thr Glu Asn Ser Leu Arg
676         435                 440                 445
678 Gly Val Ile Leu Asp Ile His Phe Leu Ser Gln Ala Asp Phe Leu Val
679     450                 455                 460
681 Cys Thr Phe Ser Ser Gln Val Cys Arg Val Ala Tyr Glu Ile Met Gln
682 465                 470                 475                 480
684 Thr Leu His Pro Asp Ala Ser Ala Asn Phe His Ser Leu Asp Asp Ile
685                 485                 490                 495
687 Tyr Tyr Phe Gly Gly Gln Asn Ala His Asn Gln Ile Ala Val Tyr Pro
688             500                 505                 510
690 His Lys Pro Arg Thr Glu Glu Glu Ile Pro Met Glu Pro Gly Asp Ile
691         515                 520                 525
693 Ile Gly Val Ala Gly Asn His Trp Asp Gly Tyr Ser Lys Gly Ile Asn
694     530                 535                 540
696 Arg Lys Leu Gly Lys Thr Gly Leu Tyr Pro Ser Tyr Lys Val Arg Glu
697 545                 550                 555                 560
699 Lys Ile Glu Thr Val Lys Tyr Pro Thr Tyr Pro Glu Ala Glu Lys
E--> 700                 565                 570                 575
2297 <210> SEQ ID NO: 64
2298 <211> LENGTH: 235
2299 <212> TYPE: PRT
2300 <213> ORGANISM: Homo sapiens
2302 <400> SEQUENCE: 64
2303 Leu Pro Ala Gln Val Ala Phe Thr Pro Tyr Ala Pro Glu Pro Gly Ser
2304   1               5                  10                  15
2306 Thr Cys Arg Leu Arg Glu Tyr Tyr Asp Gln Thr Ala Gln Met Cys Cys
2307             20                  25                  30
2309 Ser Lys Cys Ser Pro Gly Gln His Ala Lys Val Phe Cys Thr Lys Thr
2310         35                  40                  45
2312 Ser Asp Thr Val Cys Asp Ser Cys Glu Asp Ser Thr Tyr Thr Gln Leu
2313     50                  55                  60
2315 Trp Asn Trp Val Pro Glu Cys Leu Ser Cys Gly Ser Arg Cys Ser Ser
2316 65                  70                  75                  80
2318 Asp Gln Val Glu Thr Gln Ala Cys Thr Arg Glu Gln Asn Arg Ile Cys
2319                 85                  90                  95
2321 Thr Cys Arg Pro Gly Trp Tyr Cys Ala Leu Ser Lys Gln Glu Gly Cys
2322             100                 105                 110
```

RAW SEQUENCE LISTING DATE: 04/24/2006
PATENT APPLICATION: US/10/575,261 TIME: 16:15:41

Input Set : A:\seq list.txt
Output Set: N:\CRF4\04242006\J575261.raw

```
2324 Arg Leu Cys Ala Pro Leu Arg Lys Cys Arg Pro Gly Phe Gly Val Ala
2325         115                 120                 125
2327 Arg Pro Gly Thr Glu Thr Ser Asp Val Val Cys Lys Pro Cys Ala Pro
2328     130                 135                 140
2330 Gly Thr Phe Ser Asn Thr Thr Ser Ser Thr Asp Ile Cys Arg Pro His
2331 145                 150                 155                 160
2333 Gln Ile Cys Asn Val Val Ala Ile Pro Gly Asn Ala Ser Met Asp Ala
2334                 165                 170                 175
2336 Val Cys Thr Ser Thr Ser Pro Thr Arg Ser Met Ala Pro Gly Ala Val
2337             180                 185                 190
2339 His Leu Pro Gln Pro Val Ser Thr Arg Ser Gln His Thr Gln Pro Thr
2340         195                 200                 205
2342 Pro Glu Pro Ser Thr Ala Pro Ser Thr Ser Phe Leu Leu Pro Met Gly
2343     210                 215                 220
2345 Pro Ser Pro Pro Ala Glu Gly Ser Thr Gly Asp
E--> 2346 225                 230                 235
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RAW SEQUENCE LISTING ERROR SUMMARY DATE: 04/24/2006
PATENT APPLICATION: **US/10/575,261** TIME: 16:15:43

Input Set : **A:\seq list.txt**
Output Set: **N:\CRF4\04242006\J575261.raw**

Invalid Line Length:
The rules require that a line not exceed 72 characters in length. This includes spaces.

Seq#:17; Line(s) 832
Seq#:74; Line(s) 2501
Seq#:75; Line(s) 2558
Seq#:76; Line(s) 2666

VERIFICATION SUMMARY DATE: 04/24/2006
PATENT APPLICATION: **US/10/575,261** TIME: 16:15:43

Input Set : **A:\seq list.txt**
Output Set: **N:\CRF4\04242006\J575261.raw**

L:9 M:270 C: Current Application Number differs, Replaced Current Application No
L:9 M:271 C: Current Filing Date differs, Replaced Current Filing Date
L:48 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:1
L:52 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:1
L:56 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:1
L:60 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:1
L:64 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:1
L:68 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:1
L:72 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:1
L:76 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:1
L:80 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:1
L:84 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:1
L:88 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:1
L:92 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:1
L:97 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:1
L:101 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:1
L:105 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:1
L:109 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:1
L:113 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:1
L:117 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:1
L:121 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:1
L:153 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:2
M:332 Repeated in SeqNo=2
L:208 M:252 E: No. of Seq. differs, <211> LENGTH:Input:376 Found:372 SEQ:2
L:584 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:7
L:700 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:8
L:1396 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:46
L:1529 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:48
L:1676 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:51
L:1786 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:53
L:2346 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:64